

GenCore version 5.1.3
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OM nucleic - nucleic search, using sw model

Run on: February 16, 2003, 16:42:40 : Search time 2267.72 Seconds
(without alignments)
15566.316 Million cell updates/sec

Title: US-09-497-967-3
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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 24791104 seqs, 12571243825 residues
Total number of hits satisfying chosen parameters: 49582208

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Post-processing: Minimum Match 0%
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Listing first 45 summaries

Database : Pending_Patents_NA_Main:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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4	782.6	55.7	1404	18	US-09-497-967-5
5	782.6	55.7	1410	18	US-09-497-967-102
6	781	55.6	1404	18	US-09-497-967-53
7	252.6	18.0	1326	18	US-09-497-967-1
8	252.6	18.0	1326	18	US-09-498-612-7
9	252.6	18.0	2486	18	US-09-497-967-2
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11	251	17.9	2811	18	US-09-498-612-4
12	239.6	17.1	1193	3	US-07-763-352A-2
13	239.6	17.1	1193	3	US-07-763-352A-14
14	110.4	7.9	316	15	US-09-196-161-1
15	110.4	7.9	316	15	US-09-196-161-2
16	110.4	7.9	316	15	US-09-196-161-2
17	103	7.3	316	15	US-09-196-1610-9
C 18	95.2	6.8	186935	67	US-60-226-176-1976
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Sequence 75, Ap1
Sequence 71, Ap1
Sequence 7195, A
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Sequence 16038, A
Sequence 16612, A
Sequence 16768, A

RESULT 1
 US-09-497-967-3
 ; Sequence 3, Application US/09497967
 ; GENERAL INFORMATION:
 ; APPLICANT: Clark, Theodore G.
 ; APPLICANT: Dickerson, Jr., Harry W.
 ; APPLICANT: Lin, Jian-Long
 ; TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
 ; TITLE OF INVENTION: ICHTHYOPHTHIRIUS
 ; FILE REFERENCE: 235.00170101
 ; CURRENT APPLICATION NUMBER: US/09/497,967
 ; CURRENT FILING DATE: 2000-02-04
 ; PRIOR APPLICATION NUMBER: 60/131,121
 ; PRIOR FILING DATE: 1999-04-27
 ; PRIOR APPLICATION NUMBER: 60/118,634
 ; PRIOR FILING DATE: 1999-02-04
 ; PRIOR APPLICATION NUMBER: 60/122,372
 ; PRIOR FILING DATE: 1999-03-02
 ; PRIOR APPLICATION NUMBER: 60/124,905
 ; PRIOR FILING DATE: 1999-03-17
 ; NUMBER OF SEQ ID NOS: 102
 ; SOFTWARE: PatentIn Ver. 2.1
 ; SEQ ID NO 3
 ; LENGTH: 1404
 ; TYPE: DNA
 ; ORGANISM: Ichthyophthirius multifiliis
 US-09-497-967-3

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Ddb	61	GCTAATTGCCGTGGAACTGAAC	TAAACAACACCGCGATAAGTTGATGATCTAGGAAC	T 120	
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US-09-498-612-8
; Sequence 8, Application US/09498612
; APPLICANT: GAERTIG, Jacek
; APPLICANT: DICKERSON JR., Harry W.
; APPLICANT: CLARK, Theodore G.
; APPLICANT: THE UNIVERSITY OF GEORGIA RESEARCH FOUNDATION, INC
; TITLE OF INVENTION: RECOMBINANT EXPRESSION OF HETEROLOGOUS NUCLEIC ACIDS IN
; FILE REFERENCE: 235.00100101
; CURRENT APPLICATION NUMBER: US/09/498,612
; PRIOR FILING DATE: 2000-02-04
; PRIOR FILING DATE: 1999-02-04
; PRIOR FILING DATE: 1999-02-04
; PRIOR FILING DATE: 1999-03-02
; PRIOR FILING DATE: 1999-03-17
; PRIOR FILING DATE: 1999-03-17
; PRIOR FILING DATE: 1999-04-27
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; PRIOR FILING DATE: 2000-02-04
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; TYPE: DNA
; ORGANISM: Ichthyophthirius multifiliis
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; Sequence 44, Application US/09497967
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; GENERAL INFORMATION:
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; APPLICANT: Clark, Theodore G.
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; APPLICANT: Dickerson, Jr., Harry W.
;
; APPLICANT: Lin, Jian-Long
;
; TITLE OF INVENTION: DIAGNOSTIC AND
;
; TITLE OF INVENTION: ICHTHYOPHATHIRIUS
;
; FILE REFERENCE: 235,00170101
;
; CURRENT APPLICATION NUMBER: US/09497967
;
; CURRENT FILING DATE: 2000-02-04
;
; PRIOR APPLICATION NUMBER: 60/131,121
;
; PRIOR FILING DATE: 1999-04-27
;
; PRIOR APPLICATION NUMBER: 60/118,638
;
; PRIOR FILING DATE: 1999-02-04
;
; PRIOR APPLICATION NUMBER: 60/122,373
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; PRIOR FILING DATE: 1999-03-02
;
; PRIOR APPLICATION NUMBER: 60/124,900
;
; PRIOR FILING DATE: 1999-03-17
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; NUMBER OF SEQ ID NOS: 102
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; SOFTWARE: PatentIn Ver. 2.1
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; LENGTH: 1410
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; ORGANISM: Ichthyophthirius multifiliis
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; US-09-497-967-44

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RESULT 4

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; GENERAL INFORMATION:
; APPLICANT: Clark, Theodore G.
; APPLICANT: Dickerson, Jr., Harry W.
; APPLICANT: Lin, Tian-Long
; TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
; TITLE OF INVENTION: ICHTHYOPHTHIRIUS
; FILE REFERENCE: 235,00170101
; CURRENT APPLICATION NUMBER: US/09/497,967
; CURRENT FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: 60/131,121
; PRIOR FILING DATE: 1999-04-27
; PRIOR APPLICATION NUMBER: 60/118,634
; PRIOR FILING DATE: 1999-02-04
; PRIOR APPLICATION NUMBER: 60/122,372
; PRIOR FILING DATE: 1999-03-02
; PRIOR APPLICATION NUMBER: 60/124,905

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; PRIOR FILING DATE: 1999-03-17									
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; SEQ ID NO 5									
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; TYPE: DNA									
; ORGANISM: Artificial Sequence									
; FEATURE:									
; OTHER INFORMATION: Description of Artificial Sequence: synthetic									
; OTHER INFORMATION: 55kd I-antigen coding region									
US-09-497-967-5									
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Best Local Similarity 55.7%; Score 782.6; DB 18; Length 1404;									
Matches 1013; Conservative 0; Mismatches 384; Indels 0; Gaps									
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Qy	361	AATTTTATAATGAAATGCTTCCAAATTTTAAATGCAAGTGCTAGTACATGACACAGCTTGT	420						
Db	361	AACCTTCTAACAGAACGCTCCTTAACCTCAACGCTGGAGCTTCTACCTGTACCGCTTGT	420						
Qy	421	CCGGTAAACAGAGTTGGTGGTGCAATTGACTGCTGGTAAATGCCGCTACCATAGTCGCATAA	480						
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Qy	481	TGTAACGCTCCGATGCTTACTGTTGCTGAGTGTGATGATGAGTGAATCTAGTATGTT	540						
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Qy	541	AGATCATTTACAGAATGTGTTAAATGTAGACTTTTAACTTTTACTATAATGTTAATAATGGT	600						
Db	541	CGCTCTTTTACCGAGTGTGTGAAGTGTGCGCTGAACTTCTACTACACGGAACACAGGA	600						
Qy	601	AATACTCTCTTTCAATCCAGGTAAAGTTAATGCACACCTTGTTCGCGCAATTTAAACCTGCT	660						
Db	601	AACACCCCTTTCACCCCTGGAAAGTCTCAGTGTACCCCTTGTCTGCTATCAAGACCTGCT	660						
Qy	661	ATGTTGGCTTAAGCTACTTTTAGTAATGATGCTACAATAACCGCATTAATGTAACTGTTGCA	720						
Db	661	AACGTGGGTCTAGGCTAGCCCTGGGAAACGACGCTACCATCACCGCTCAGTGTACGTGGCT	720						
Qy	721	TGCGCTGATGTTACTATAAGTGTCTGAGTGAATTAATATTTGGGTAGCAACAAACACTGAA	780						
Db	721	TGTCCTGAGGAACCATCTCTGCTGTGGAGTGAACAACACTGGGTGGCTCAGAACACCCGAG	780						
Qy	781	TGTACTAATTTGCTCCTTAACCTTTTACAATAATAATGCTCCTTAATTTCAATCCAGTAA	840						
Db	781	TGTACCAACTGTGCTCCTAACTTCTACACAAACACAGCTCTTAATCTTCAACCCCTGGAAAC	840						
Qy	841	AGTACATGCCTACCTTGCCTCCACGAATAAAGATTAATGTTGCTGAAGCCACTGCAGGTGCT	900						

Matches 1013; Conservative		0; Mismatches	384; Indels	0; Gaps
Qy	1	ATGAAAATAATATTTT	AGTAATATGATTATTT	CAATATTTATTTATCAATTAATAATCT 60
Db	1	ATGAAGAACACATCCT	GGTGATCCTGATCNCT	CTCTGTTCATCACCAGATCAAGTCT 60
Qy	61	GCTAATTGTCCTGTTG	GAACTGAACACACCGCGGAT	TAAGTTGATGATCTAGGAATCT 120
Db	61	GCTAACTGTCCTGTTG	GAAACGAGAACACACCGCTGGACAGG	TGGACGACCTTGGAAACC 120
Qy	121	COTGCAAAATGGTTAA	TTGTTAGAAAACCTTTT	ATTAATATGATGCTGCTTTCGTT 180
Db	121	CCTGCTAACTGTGTGA	AGTGTCAAGAAGATTT	CTACTACAAACGCTGCTGTTTCGTG 180
Qy	181	CCTGGTGTAGTACGT	GATACACCTTGTCCATAAAAA	AGATGCTGTGTTTAAACCAAT 240
Db	181	CCTGGAGCTTCTACCT	GATGACCCCTTGCTCAGAA	GAGGACGCTGAGCTCAGCCTAAC 240
Qy	241	CCACCTGCTACGTGCT	TAATTTAGTACATAA	TGTRACGTTAAATGCCCTGCTGTCACGCA 300
Db	241	CCTCCTGCTACCGCT	TAACCTGGTGACCCAGGT	TACGTTGAAGTGCTCTGCTGGNACCGCT 300
Qy	301	ATTGCAAGTGGAGCA	ACAGATTATCAGCAAT	TAATCAGAGAATGTGTTAATTTGTAAGTT 360
Db	301	ATCGCTGGAGGACCT	TACCGACTACGCTGCT	ATCATCACCGAGTGTGAACTGTGCGCATC 360
Qy	361	AATTTTATAATGA	AAATGCTCCAAATTTT	TAATGCAGTGTCTAGTACATGCACAGCTTGT 420
Db	361	AACCTTCTACAAC	GAGAACCTCCTCAACT	CTCAACGCTGGAGCTTCTACTGTACCGCTTGT 420
Qy	421	CCGGTAAACAGAGT	CTGGTGTCAATGCTG	TGTAATGCCGTACCATAGTCGCAATAA 480
Db	421	CCGTGTAAACCGCT	GGGAGAGCTTGACCGCT	GGAACGCTGCTACCATCGTGGCTCAG 480
Qy	481	TGTAACGCTGCATG	CTCTACTGCTACTGC	ATGCATTTGATGAGTAGTACTTATGTT 540
Db	481	TGTAACGCTGCTTGT	CTCCACGGAACCGCTCT	GGAACGAGTGACCCAGCTACGCTG 540
Qy	541	AGATCATTCACAA	TGTGTTAAATGTACATTA	ACTTTTACTATAATGTTAATAATGGT 600
Db	541	CGCTCTTTACCGAG	TGTGTGAAGTGTCGCC	TGTAAGTGTGCCCTGTAACCTACTACAACGGAACACGGA 600
Qy	601	AATACTCCTTTCA	ATCCAGTAAAGTTAAT	GCACACCTTTGCCGGCAATTAACCTGTCT 660
Db	601	AACACCCCTTTCA	ACCTGAAAGTCTCAGT	GTACCCCTTGTCCTGCTATCAAGCTGCT 660
Qy	661	AATGTGCTTAAG	CTACTTTAGTAAATG	TATGATGCATAACCGCATATGTAACGTTGCA 720
Db	661	AACGTGCTCAGG	CTACCTTGGAAACGAGCT	TACCATCACCCTCTCAGTGTAACTGGCT 720
Qy	721	TGCTGATGCTACT	ATAAGTGTCTGGAGTA	AAATTAATTTGGGTAGCCACAAACACTGAA 780
Db	721	TGCTCTGACGA	ACCATCTCTGCTGGAGT	GAAACAACCTGGGTGGCTCAGAACACCGAG 780
Qy	781	TGTACTAATTTG	CTCTACTTTTACAATA	TAAATGCTCAATTTCAATCCAGTAAAT 840
Db	781	TGTACCACTGTG	CTCTAACTTCTACAACA	CAAGCTCCTTAACCTTCAACCTCGAAAC 840
Qy	841	AGTACATGCC	TTFACCTTGCACGAA	ATAAGATATATGGTGTGCTGAAGCCACTGCAGGTGGT 900
Db	841	TCTACTGTCTG	CTCTGCTTGTCTTACA	AGGACTACGGAGCTGAGGCTACCGCTGGAGA 900
Qy	901	CGCGCTACTTTAG	CCAAATAAATGTAAT	TATTTGATGCCCTGATGGTACTGCAATTTGCTAGT 960
Db	901	GCTGCTACCC	TGGGTGAAGCTGTAA	ATCATCGCTTGTCTGACGGAACCGCTATCGCTTCT 960
Qy	961	GGAGCAACTAAT	TATGTAAATATAACA	AGAATGTCTAAATTTGCTGCTCACTTTTAT 1020
Db	961	GGAGCTACCACT	ACTAGCTGATCCTCC	ACACCGAGTGTCTGAATGTCTGCTAACTTCTAC 1020
Qy	1021	TTTGTGTTAATA	TTTCTAGGCAAG	AAAGTAGTAGTGAACACCATGTCCACCAATAAA 1080
Db	1021	TTTCAGCGGA	AAACAACTTCCAGGCT	GGATCTCTCGCTGTAAAGCTTGTCTCTGCAACAAG 1080

QY	1081	GTTTAAGCGCTAGCAACTGCAGGTGCTACTTCTTAAATTGCATAATGTGCCCTT	1140
Db	1081	GTGAGGAGCTGTGGCTACCGTGGAGAACCGTACCTCATCGCTCAGTGTGCTGTG	1140
QY	1141	GAATGCCCTGCTGGTACTGTACTCACCGATGAACAACATCTACTTATAATAGCAGCA	1200
Db	1141	GAGTGTCTCTGTAACCGTGTGACCGACGGAACACCTCTACCTACAAGCAGGCTGCT	1200
QY	1201	TCTGAATGTGTTAAATGTGTCGCCACTTTTATATCTACAAAATAAAGCTGATTGGGTAGCA	1260
Db	1201	TCTGAGTGTGTAAGTGTGCTGTGTAATTTTCTTACACCAACCAACGACCTGGGTGCTG	1260
QY	1261	GGTATTGATCATGTACTACTTCTTAATAAAAAATTAACCTTCTGGCGCTGAAGCTAAATTTA	1320
Db	1261	GGATTCACACCTGTACCTCTTGTAAACAAGAAGCTGACCTCTGGAGCTGAGGCTAACCTG	1320
QY	1321	CCTGAATCTGCTAAAAAATATATATGTGATTTGCCTAAATTTTATCAATTTCTCTTA	1380
Db	1321	CCTGAGTCTGCTAAGAAGACATCCAGTGTGACTTCGCTAACTTCTGTCTATCTCTCTG	1380
QY	1381	TTATTGATTTCTTTATTA	1397
Db	1381	CTGTGATCTCTTACTA	1397
RESULT 6			
US-09-497-967-53			
; Sequence 53, Application US/09497967			
; GENERAL INFORMATION:			
; APPLICANT: Clark, Theodore G.			
; APPLICANT: Dickerson, Jr., Harry W.			
; APPLICANT: Lin, Tian-Long			
; TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF			
; TITLE OF INVENTION: ICHTHYOPHIRIUS			
; FILE REFERENCE: 235.00170101			
; CURRENT APPLICATION NUMBER: US/09/497,967			
; CURRENT FILING DATE: 2000-02-04			
; PRIOR APPLICATION NUMBER: 60/131,121			
; PRIOR FILING DATE: 1999-04-27			
; PRIOR APPLICATION NUMBER: 60/118,634			
; PRIOR FILING DATE: 1999-02-04			
; PRIOR APPLICATION NUMBER: 60/122,372			
; PRIOR FILING DATE: 1999-03-02			
; PRIOR APPLICATION NUMBER: 60/124,905			
; PRIOR FILING DATE: 1999-03-17			
; NUMBER OF SEQ ID NOS: 102			
; SOFTWARE: PatentIn Ver. 2.1			
; SEQ ID NO 53			
; LENGTH: 1404			
; TYPE: DNA			
; ORGANISM: Artificial Sequence			
; FEATURE:			
; OTHER INFORMATION: Description of Artificial Sequence: synthetic G5			
; OTHER INFORMATION: proline mutant i-antigen			
US-09-497-967-53			

Query Match	55.6%	Score 781;	DB 18;	Length 1404;
Best Local Similarity	72.4%	Pred. NO. 2.9e-174;		
Matches 1012: Conservative	0;	Mismatches 385;	Indels 0;	Gaps 0;

[illegible]

QY 181 CCTGGTCTAGTACGTGTACACCTGTGCCATAAAAAAGATGCTGGTCTTAACCAAT 240
DB 181 CCTGGAGCTTCTACCTGTACCCCTGTCTCGAAGAAAGAGCGCTGGAGCTCAGCGCTAAC 240
QY 241 CCACCTGCTACTGCTAATTTAGTACATATGTAAAGTTTAAATGCCCTGCTGGTACCGCA 300
DB 241 CCTCCTGCTACCGCTAACCTGTGTGACCCAGTGTAAAGTGAAGTGTCTGCTGGAACCGCT 300
QY 301 ATTGCAGGTGGAGCAACAGATTATGACGAGTAATACACAGAATGTGTTAATGTAGAATT 360
DB 301 ATGCTGGAGGAGCTACCGACTACGCTGTATCATCACCAGTGTGTGAACCTGTGCGATC 360
QY 361 AATTTTATAATGAAATGCTCCAAATTTAATGCAAGTGTGAGTACATGACACAGCTTGT 420
DB 361 AACTTCTACAACGAGAAGCTCTTACTTCAACGCTGAGCTTCTACCTGTACCGCTTGT 420
QY 421 CCGGTAAACAGATTGGTGGTGCATGTGCTGTGTAATGCGGCTPACCATAGTGCATAA 480
DB 421 CTTGTGAACCGTGTGGGAGGAGCTGTGACCGCTGGAACCGCTGTACCATCGTGCCTAG 480
QY 481 TGTAACTTACAGAAATGTTTAAATGTAGACTTAACTTTTACTATATGTTAATGTTAATGTT 600
DB 541 CGCTCTTTTCAACCGAGTGTGAAGTGTGCGCTGAACTTCTACTACAACGGAACACGGA 600
QY 601 AATACTCTCTTCAATCCAGGTAAGTTTAAATGACACACCTTGTGCGGCAATTAACCTGCT 660
DB 601 AACACCCCTTTCAACCCCTGGAAAGTCTCAGTGTACCCCTTGTCTGTCTATCAAGCTGCT 660
QY 661 AATGTTGCTTAAGTACTTAACTTAACTGATGCTACAAATACCGCATATGTTAAGTTGCA 720
DB 661 AACGTGCTGCTGAGCTACCTGCGGAAACGACGCTACCATCACCGCTCAGTGTAAAGTGGCT 720
QY 721 TGCCCTGATGTPACTATAGTGTCTGCTGGAGTAATTAATGGGTAGCACAAAACACTGAA 780
DB 721 TGCTCTGAGGAAACCATCTGCTGCTGGAGTGAACAACTGGTGGCTCAGAACACCGAG 780
QY 781 TGTACTAATTTGCTCTCTCACTTTTACAATAATAATGCTCTCTAATTTCAATCCAGTTAAT 840
DB 781 TGTACCAACTGTGCTCTCACTTCTACACAAACACGCTCTTACTTTCAACCCCTGGAAC 840
QY 841 AGTACATGCTTACCTTGGCCAGCAAAATAAGATTATGCTGTGAAGCACTGCGAGTGGT 900
DB 841 TCTACCTCTGCTGCTTGTCTGCTTAACAAGGACTACGGAGCTGAGGCTACCGCTGGAGGA 900
QY 901 GCGGCTACTTTAGCAATAATGTAATTTGCATGCGCCCTGATGGTACTGCAATTTGCTAGT 960
DB 901 GCTGCTACCTTGGCTAAGCAGTGTAAACATCGCTTGTGCTGACGGAACCGCTATCGCTTCT 960
QY 961 GGAGCACTAATATGTAATATTATAACAGAATGTCTAAATTTGCTGCTACTCTTTAT 1020
DB 961 GGAGCTACCAACTACGTGATCTCTGACAGCCGAGTGTCTGACTGTGCTGCTAATCTTCTAC 1020
QY 1021 TTTGATGTAATATTTCTAGGAGGAGTGTAGTAGTGAAGCAAGCTGTCCAGCAATAAA 1080
DB 1021 TTCACGGAACAACACTTCCAGGCTGGATCTTCTCGTGTAAAGCTTGTCTGCTGAACAAG 1080
QY 1081 GTTTAAGCCGCTGTAGCAACTGCAAGTGTGCTACTTCTAATTTGCTAATGTTGCCCTT 1140
DB 1081 GTGCAGGAGCTGTGGCTACCGCTGGAGGAGCCGCTACCCGTATGCTCAGTGTGCTCTG 1140
QY 1141 GAATGCCCTGCTGTACTGTACTCAGGATGGAAACAACTACTTATATAAATGAACGCA 1200
DB 1141 GAGTGTCTGCTGGAACCGCTGTGACCGAGCAACCCCTACCTTACAGCAGGCTGCT 1200
QY 1201 TCTGAATGCTGTTAAATGTGCTGCCAACTTTTATACTACAAATAAAGTATTTGGGTAGCA 1260
DB 1201 TCTGAGTGTGAGTGTGCTGCTAACTTCTACACCAACCAACGACGACCTGGGTGCT 1260
QY 1261 GGTATTGATACATGTACTAGTTGTATATAAAAAAATAAATCTCTGCGGCTGAAGCTAATTA 1320

DB 1261 GGAATCGACACCTGTACCTTGTACACAGAAGCTGACCTCTGGAGCTGAGGCTAACCTG 1320
QY 1321 CCTGAATCTGCTAAAAAATAATATATGTGATTTGGTAAATTTTTTATCAATTTTCTTAA 1380
DB 1321 CCTGAGTCTGCTAAGAAGAACAATCCAGTGTGACTTTCGCTAACTTCTGCTATCTCTCTG 1380
QY 1381 TTATTGATTTCTTTATTA 1397
DB 1381 CTGCTGATCTCTTACTA 1397

RESULT 7

US-09-497-967-1
; Sequence 1, Application US/09497967
; GENERAL INFORMATION:
; APPLICANT: Clark, Theodore G.
; APPLICANT: Dickerson, Jr., Harry W.
; APPLICANT: Lin, Tian-Long
; TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
; TITLE OF INVENTION: ICHTHYOPHTHIRIUS
; FILE REFERENCE: 235, 00170101
; CURRENT APPLICATION NUMBER: US/09/497,967
; CURRENT FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: 60/131,121
; PRIOR FILING DATE: 1999-04-27
; PRIOR APPLICATION NUMBER: 60/118,634
; PRIOR FILING DATE: 1999-02-04
; PRIOR APPLICATION NUMBER: 60/122,372
; PRIOR FILING DATE: 1999-03-02
; PRIOR APPLICATION NUMBER: 60/124,905
; PRIOR FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 102
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 1326
; TYPE: DNA
; ORGANISM: Ichthyophthirius multifiliis
US-09-497-967-1

Query Match 18.0%; Score 252.6; DB 18; Length 1326;
Best Local Similarity 56.7%; Pred. No. 5.2e-49;
Matches 660; Conservative 0; Mismatches 394; Indels 111; Gaps 6;

QY 344 GTGTTAATTTGTAAGTAATTTTATTAATGAAATGCTCCAAATTTTAAATGCAAGTGTCTA 403
DB 167 GTGCTGCTTAAAGAGAAAGCTAATGGTAATTAACCTTTTCGACGCAAAATATGCTGTAGAG 226
QY 404 GTACATGCACAGCTTGTCCGTAACAGAGTTGGTGGTGCATTTGACTGCTGGTAAATGCCG 463
DB 227 GTATATGTGTACCATGCCAATAAAGAGATGAGTCTGTGTACCAATGCAAGTGTAGTAG 286
QY 464 CTACCATAGTGCATAATGTAAGTGCATGCTCTACTGCTGCTACTGCTGCTGCTGATGGAG 523
DB 287 CTACTTTAGCCACATAATATGCTGCTACTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 346
QY 524 TAACACTGCTATTTAGTATGATCATTCACAGAATGTTTAAATGTAGACTTAACTTTTACT 583
DB 347 TGACAGATGTTTTGATAGATCAGCCGCAATAATGTGTTAAATGCAAAACCTTAATTTACT 406
QY 584 ATATGTAATAATGTTAATGTAATCTCTTCAATCCAGTAAAGTTAATGCAACCTTGTCTG 643
DB 407 ATATGTTGTTCTCTCTTAAAGTGAAGCTTCCGCTGCTGCTGCTGCTGCTGCTGCTGCTG 466
QY 644 CGGCAATTAACCTGCT-----AATG 664
DB 467 CCGCTGCAAGTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 526
QY 665 TTGCTTAAGCTACTTTAGGTAAATGATGCTACATAACCCATTAATGTAACGTTGATGCC 724
DB 527 ATTCTCTCCCTGCACTGCGAGTGCCTAAGCTAAATTTAGCCACATAATGTAGCAATTAATGTC 586
QY 725 CTGATGCTACTATAGTGTGCTGCTGGAGT---AAATAATTTGGGTAGCACAACAACTGAAT 781

Thu Feb 20 11:10:17 2003

Db 587 CTACTGGCAGCTGTTACTTGTATGATGAGTGACACTGTGTTTAAATACATCAGCCACATTAAT 646
QY 782 GTACTAATGTCCTTAACATTTTACAAATAATAATGCTCCTCAAT 826
Db 647 GTGTTAAATGACAGACCTAATCTTTACTTATATGATGCTGCTTAAAGGTGAAGCTCCTG 706
QY 827 -----TCAATCCAGGTAAATAGTACAT 847
Db 707 GCGTTTAAAGTTTTTGTGCTGCTGCTGCCGCTGCAGAGTGTGCTGCCGTTACTAGTTAAT 766
QY 848 GCGTACCTTGCCAGCAAAATAAAGATTATGCTGAAGCACTGCAGGTGGTCCGCTA 907
Db 767 GTGTACCTTGCCAAATAAACAARAACGATTTCTCT---GCCACTGCAGGTGCCTAAGCTA 823
QY 908 CTTTAGCCAAATAATGTAATATGTCAGCCCTGATGCTACTGCTAATGCTAGTGGAGCAA 967
Db 824 ATTTAGCCACATAATGCAGTACTTAATGTCCAAGTGGCACTGCAATTTCAAGACGGAGTGA 883
QY 968 CTAATTAATGTAATATATAACAGAGATGCTAAATTTGCTGCTAACTTTTATTTTGATG 1027
Db 884 CACTTGTGTTTAAATCATCCACATAATGTTCTTAATGCAATGCTAAATTTTAA 943
QY 1028 GTAATAATTTCTAGCAGGAAGTACTAGATGCAAGCACTGCCAGCAATAAAGTTTAAAG 1087
Db 944 ATGGTAATTTCCAGCAGAGTAAAGTTAATGTTTAAAGTGTCCAGTAAGTAAACT---A 1000
QY 1088 GCGCTGTAGCAACTGCAGGTGGTACTGCTACTTTAATTTGATGATGCTGCTGATGCTG 1147
Db 1001 CTCAGCACATGCTCCAGGTAATGCTACTTAAAGCCACACATAATGTTTACCACATGTC 1060
QY 1148 CTGCTGGTACTGACTCACCGATGAACAACATCTACTTATAAATAAGCAGCATCTGAAT 1207
Db 1061 CTGCTGGTACAGTACTGATGATGAACATCAACTTAATTTGTAGCTTCCGCACTGAAT 1120
QY 1208 GTGTTAAATGCTGCGCAACTTTTATCTACAAATAAATAAATGATGCTGCTAGCAGGTATG 1267
Db 1121 GTACTAATGTTCTGCTGCTTTTGTGATCAAAAAACAACCTGGTTTACAGCAGGTACTG 1180
QY 1268 ATACATGCTAGTCTGTAATAAATAAATAAATCTGCGCTGAAGCTAATTTACTCTGAAT 1327
Db 1181 ATACATGCTAGTGAATGACTAAAAAATAAATCTGCTGCTGCTGCTGCTGCTGCTGCTG 1240
QY 1328 CTGCTAAAAAATAATATAATG-----TCATTTGCTCAATTTTATCAATTTCCCT 1378
Db 1241 AAGCTACTCAAAAGTATATATGCGCTCCACTACTTTCGCTAAATTTTATCGATTTCCCT 1300
QY 1379 TATTATTGATTTCTTATTATTATT 1403
Db 1301 TATTATTATTCTTCTATTATT 1325

RESULT 8
US-09-498-612-7
; Sequence 7, Application US/09498612
; GENERAL INFORMATION:
; APPLICANT: GAERTIG, Jacek
; APPLICANT: DICKERSON Jr., Harry W.
; APPLICANT: CLARK, Theodore G.
; APPLICANT: THE UNIVERSITY OF GEORGIA RESEARCH FOUNDATION, INC
; TITLE OF INVENTION: RECOMBINANT EXPRESSION OF HETEROLOGOUS NUCLEIC ACIDS IN
; FILE REFERENCE: 235.00100101
; CURRENT FILING DATE: 2000-02-04
; PRIOR FILING DATE: 60/118,634
; PRIOR FILING DATE: 1999-02-04
; PRIOR FILING DATE: 60/122,372
; PRIOR FILING DATE: 1999-03-02
; PRIOR FILING DATE: 60/124,905
; PRIOR FILING DATE: 1999-03-17
; PRIOR FILING DATE: 60/131,121
; PRIOR FILING DATE: 1999-04-27

; PRIOR APPLICATION NUMBER: PCT/US00/02966
; PRIOR FILING DATE: 2000-02-04
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 7
; LENGTH: 1326
; TYPE: DNA
; ORGANISM: Ichthyophthirius multifiliis
US-09-498-612-7

Query Match 18.0%; Score 252.6; DB 18; Length 1326;
Best Local Similarity 56.7%; Pred. No. 5.2e-49;
Matches 660; Conservative 0; Mismatches 394; Indels 111; Gaps 6;

QY 344 GTGTTAATTTGTAGAATTAATTTTATATGAAATGCTCCAAATTTTAAATGCAAGGTGCTA 403
Db 167 GTGCTGCTTAAAGGAGAAGCTAATGGTAATTAACCTTTCCGACGAAATAATGCTGCTAGAG 226
QY 404 GTACATGACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACGTGGTAAATGCCG 463
Db 227 GTATATGTTACCATGCCAAATAAACAGAGTAGGCTCTGTTACCAATGCAAGTGAAGTCTAG 286
QY 464 CTACCATAGTGCATAATGTTAACGTGCGCATGCTCTACTGCTACTGCTACTGCTACTGATGGAG 523
Db 287 CTACTTTAGCCACATAATGCGACTACTTAATGCTCTACTGCGCACTGCACTTGTATGGAG 346
QY 524 TAACACTGATTATGTTAGATCATTCACAGAAATGTTTAAATGTAGACTTAACCTTTTACT 583
Db 347 TGACAGATGTTTTTGTAGATCAGCCGCAATATGTTTAAATGCAACCTTAACCTTTTACT 406
QY 584 ATAATGGTAATGTTAATGTTAATCTCTCTTTCAATCCAGGTAAAGTTAATGCAACCTTTGTC 643
Db 407 ATAATGGTGGTTCTCTTAAAGTGAAGCTCTGCGGTTTAAAGTTTTCGCTGCTGCTG 466
QY 644 CGCAATTAACCTGCT-----AATG 664
Db 467 CCGCTGCAGGTGTGCTGCCGTTACTAGTTAATGTTACTTTGCCAATAAACAACG 526
QY 665 TTGCTTAAAGCTACTTTAGTAAATGATGCTACATAACCGCATATGTAACCTTGCATGCC 724
Db 527 ATTCTCTGCCACTGCGAGGTGCTAAGCTAATTTAGCCACATAATGTAGCAATTAATGTC 586
QY 725 CTGATGCTACTATAAGTGTGCTGGAGT---AAATAATGGGTAGCAGCAAAACACTGAAT 781
Db 587 CTACTGCACTGTACTTATGATGATGGAGTGACACTTGTGTTTAAATACATCAGCCACTTAT 646
QY 782 GTACTAATTTGCTCTTAACCTTTTACATAATAATGCTCCTAAT----- 826
Db 647 GTGTTAAATGCAACCTTAACCTTTTACTATAATGGTGGTTCCTCTTAAGGTGAAGTCTCTG 706
QY 827 -----TCAATCCAGGTAAATAGTACAT 847
Db 707 GCGTTTAAAGTTTTTGTGCTGCTGCCGCTGCAGGTGTGCTGCCGTTACTAGTTAAT 766
QY 848 GCGTACCTTGCCAGCAAAATAAAGATTATGCTGGAAGCCACTGCAGGTGGTCCGCTA 907
Db 767 GTGTACCTTGCCAAATAAACAARAACGATTTCTCT---GCCACTGCAGGTGCTTAAAGTGA 823
QY 908 CTTTAGCCAAATAATGTAATTTTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 967
Db 824 ATTTAGCCACATAATGCAAGTACTTAATGTTCCAACTGGCACTGCAATTTCAAGACGGAGTGA 883
QY 968 CTAATTAATGTAATATATAACAGAGATGCTAAATTTGCTGCTGCTAATTTTATTTTATG 1027
Db 884 CACTTGTGTTTAAATTCATCCACATAATGTTCTTAAATGCAATGCTAATTTACTTTTAA 943
QY 1028 GTAATAATTTCTAGCAGGAAGTACTAGATGCAAGCACTGCCAGCAATAAAGTTTAAAG 1087
Db 944 ATGGTAATTTCCAGCAGAGTAAAGTTAATGTTTAAAGTGTCCAGTAAAGTAACT---A 1000
QY 1088 GCGCTGTAGCAACTGCAGGTGGTACTGCTACTTTAATTTGATGATGCTGCTGCTGCTGCTGCT 1147
Db 1001 CTCAGCACATGCTCCAGGTAATGCTACTTAAAGCCACATAATGTTTACCACATGTC 1060

QY	644	CGCAATTTAACTGGT-----AATG	664
Db	899	COCGTCAGGTGTTCGCGTTACTAGTTAATGTGTACCTTGCCAATAACAANAACG	958
QY	665	TTGCTTTAAAGCTACTTTAGTAGTAATGATCTACAATAACGCATATGTAAAGTTGCGATGCC	724
Db	959	ATTCTCCTGCCACTCGAGTGCCTAAGCTAATTTAGCCACATATGTAGCAAATTAATGTGTC	101
QY	725	CTGATGGTACTAATAGTCTGCTGGAGT---AAATAATTTGGGTAGGACAAAACACTGAAT	781
Db	1019	CTACTGGCACTGACTTCTGATGAGGTGACACTTGTTTTTAATATACATCAGCCACATTAAT	107
QY	782	GTAATAATTGTGCTCCTAACCTTTTACAATAATAATGCTCCCTAATT-----826	
Db	1079	GTGTTTAAATGCAGACCTAACTTTTACTATATAGTGTGTTCTCCTTTAAGGTGAAGCTCCG	1139
QY	827	-----TCATCCAGGTAATAGTACAT	847
Db	1139	GGGTTTAAGTTTTGCTGCTGGTCTGCCGCTCGAGGTGTGCTGCGCTACTAGTAAAT	1199

[illegible]

Qy	1268	ATACGTGCTACTAGTTCCTGAATAAAAAATTAACTTCTGGCGCTGAAGCTAAATTTACCTGAAT	1327
Dd	1613		
Qy	1328	CTGCTAAAANAATATATAATG-----TGATTTGCCTAAATTTTTTATCAATTTCCCT	1672
Dd	1673	AGCTACTACAAAAGTATAATCGCGCTCCACTACTTTCGCTAAATTTTTTATCGATTTCCCT	1732
Qy	1379	TATTATTGATTTCTTATTATTTATT	1403
Dd	1733	TATTATTATTCTTCTTATTATT	1757

RESULT 10
US-09-498-612-3
; Sequence 3, Application US/09498612
; GENERAL INFORMATION:
; APPLICANT: GNERTIG, Jacek
; APPLICANT: DICKERSON JR., Harry W.
; APPLICANT: CLARK, Theodore G.
; TITLE OF INVENTION: RECOMBINANT EXPRESSION OF HETEROLOGOUS NUCLEIC ACIDS
; TITLE OF INVENTION: PROTOZOA
FILE REFERENCE 035 001000


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Db 1649 GTGCTGTTAAGGAGAAAGCTAAATGGTAATTAACCTTTCGACGCAATATAGCTGCTAGAG 1590
Qy 404 CTACATGCACAGCTTGTCGGTAAACAGAGCTTGGTGGTGCATAGCTGCTGGTAATGCCG 463
Db 1589 GTATATGTGTACCAATGCAAGAGAGTGGCTGTCTGTACCAATGCAGGTGACTTAG 1530
Qy 464 CTACCATGTGCGCAATATGTAAAGCTGCGCATGTCTACTGTGCTGCATCTGTATGATGGAG 523
Db 1529 CTACTTTAGCCACATATAGCAGTACTTAATGTCTTACTGGCAGCTGCATGTATGATGGAG 1470
Qy 524 TAACCTACTGATATGTAGATCAATTCACAGATGTGTTAAATGTAGACTTAACCTTTTACT 583
Db 1469 TGACAGATGTTTGTAGATAGTACGCCGCAATATGTGTTAAATGCAAACTAACTTTTACT 1410
Qy 584 ATAATGTAATAATGTGTAATACTCTTCAATCCAGGTAAAGTAAATGTCACACACTTGTG 643
Db 1409 ATAATGTTGTTCTCTTAAGGTGAAGCTTCCCTGGCGTTTAACTTTTGTGCTGCTGCTG 1350
Qy 644 CGCAATTAACCTGCT-----AATG 664
Db 1349 CCGCTGAGGTGTGCTGCGCTTACTAGTTAATGTACTTGTCCCACTAAACAAACG 1290
Qy 665 TTGCTTAAAGCTACTTTAGGTATATGCTTACAAATACCGCATATATGTAACTGTCATGCC 724
Db 1289 ATTCTCTGCCACTGCAGTGCGCTAAGCTAATTTAGCCACATATATGTAGCAATTAATGTC 1230
Qy 725 CTGATGGTACTATAAGTGTGCTGCTGAGT---AAATAATTTGGGTAGCACAACAACTGAAT 781
Db 1229 CTACTGCCACTGTACTTGTATGATGAGTGAGTGACACTTGTGTTTAAATACATCAGCCACAT 1170
Qy 782 GTACTAATTTGCTGCTTAACTTTTACAATAATAATGCTCCTAATTT-----826
Db 1169 GTGTTAATGCAGACCTAACTTTTACTATAATGGTGGTGTCTCCTTAAGGTGAAGCTCCTG 1110
Qy 827 -----TCAATCCAGGTAAATAGTACAT 847
Db 1109 CGGTTTAAGTTTTTCTGCTGCTGCGGTGAGGTGTGCTGCGGTACTAGTTAAT 1050
Qy 848 GCTACCTTGCCACACAATAAAGATATATGCTGCTGAAGCCACTGCAGGTGGTGGCCGCTA 907
Db 1049 GTGTACCTTGCCAAATAAACAACAAACGATTTCTCT---GCTCTGCAGGTGCCTAAGCTA 993
Qy 908 CTTTAGCCAAATATGTAATATTCATGCTGCTGATGCTGCTGCTGCTGCTGCTGCTGCTGCT 967
Db 992 ATTTAGCCACATATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 933
Qy 968 CTAATATGTATATATAAACAAGATGCTTAAATTTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1027
Db 932 CACTTGTGTTTGTAGTAATTCATCCACATATGTTCTTAATGCATGCTAATTTTATTTTA 873
Qy 1028 GTAATAATTTCTAGCCAGGAGTAGTAGATGCAAGCATGCTCAGCAAAATAAAGTTTAAAG 1087
Db 872 ATGGTAATTTTGAAGCAGGTAAGTAAATGTTTAAAGTGTCCAGTAAGTAAACT---A 816
Qy 1088 GCGTGTAGCAACTGCAGGTGGTACTGCTACTTTTAAATTTGATATATGCTGCTGCTGCTGCT 1147
Db 815 CTCAGCACATGCTCCAGGTAATCTGCTACTTAAGCCACATATGTTTGTAGCTTCCGCAACTGA 756
Qy 1148 CTGCTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1207
Db 755 CTGCTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 696
Qy 1208 GTGTTAAATGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1267
Db 695 GTACTAAATGTTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 636
Qy 1268 ATACATGCTAGTTGTAATAAATAATTAACCTTCTGCGCTGAAGCTAATTTTACCTGTAAT 1327
Db 635 ATACATGCTAGTTGTAATAAATAATTAACCTTCTGCGCTGAAGCTAATTTTACCTGTAAT 576
Qy 1328 CTGCTAAATAAATAATATATATG-----TGATTTCTGCTAATTTTATCAATTTCTCT 1378
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Db 575 AAGTACTCAAAAAGTATAATGCGCCTCCACTACTTTTCGCTAAATTTTATCGATTTCCT 516
Qy 1379 TATTATTGATTCTCTTATTATTATTAT 1403
Db 515 TATTATTATTCTTCTCTATTATTAT 491
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RESULT 12
US-07-763-352A-2
; Sequence 2, Application US/07763352A
; GENERAL INFORMATION:
; APPLICANT: Clark, Theodore G.
; APPLICANT: Dickerson, Harry W.
; TITLE OF INVENTION: ICH IMMOBILIZATION ANTIGEN AND FISH
; TITLE OF INVENTION: VACCINE
; NUMBER OF SEQUENCES: 15
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Greenlee and Winner
; STREET: 5370 Manhattan Circle, Ste. 201
; CITY: Boulder
; STATE: Colorado
; COUNTRY: USA
; ZIP: 80303
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07763,352A
; FILING DATE: 19910920
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Ferber, Donna M.
; REGISTRATION NUMBER: 33,878
; REFERENCE/DOCKET NUMBER: 15-91
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 303/499-8080
; TELEFAX: 303/499-8089
; TELEX: 823189
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1193 base pairs
; TYPE: NUCLEIC ACID
; STRANDEDNESS: double
; TOPOLOGY: linear
; MOLECULE TYPE: cdna to mRNA
; FEATURE:
; NAME/KEY: mat_peptide
; LOCATION: 13..1125
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 1..1128
; OTHER INFORMATION: /codon= (seq: "taa", aa: Gln)
; OTHER INFORMATION: /codon= (seq: "tag", aa: Gln)
; US-07-763-352A-2
```

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Query Match 17.1%; Score 239.6; DB 3: Length 1193;
Best Local Similarity 55.2%; Pred. No. 6e-46;
Matches 551; Conservative 0; Mismatches 414; Indels 33; Gaps 3;
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Qy 344 GTGTTAATGTGAGAATTAATTTTATAAATGCTCCAAATTTTATCGAGGTGCTA 403
Db 119 GTGCTGTTAAGGAGAGCTAATGGTAATTAACCTTTCGACGCAATATAGTGTAGAG 178
Qy 404 GTACATCCACAGCTGTGCGGTAAACAGAGTTGGTGGTGCATTTGACTGCTGTAATGCCG 463
Db 179 GTATATGTGTACCATGCCAAATAAACAGAGTAGGCTCTGTTACCAATGCAGGTGACTTAG 238
Qy 464 CTACCATAGTCGCAATATGTAACGTCGCATGCTCTACTGCTACTGCATTTGATGGAG 523
Db 239 CTACTTTAGCCACATATATGCAGTACTTAATGCTTCTACTGCGCACTGCATTTGATGGAG 298
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QY 524 TAACTACTGATTATGTTAGATCAATTCACAGAAATGTTAAATGATGAGACTTAACCTTTTACT 583
Db 299 TGCAGATGTTTTTGATAGATCAGCGCATATGTTGTTAAATGCAAACTAACTTTTACT 358
QY 584 ATAATGGT-----AATAATGGTAAATACCTTTCAATC 616
Db 359 ATAATGGTGTCTCTCTTAAGGTGAAGCTCCTGGCGTTTAAAGTTTTCGCTGCTGGTCTG 418
QY 617 CAGGTAAGATTAAGTCACACCTTGTCGGCAATTAACCTGCTAAATGTTGCTTTAAGCTA 676
Db 419 CCCTCAGGTGCTAAGCTAAATTTAGCCACATAATGTAGCAATTAATGCTTACTGGCA 478
QY 677 CTTTAGGTAATGCTACATAAACCAGCAATATGTAAGCTGTCATGCCCTCATGGTACTA 736
Db 479 CTGACTTGATGATGAGTGACACTGTTTAAATACATCAGCCACATTAATGTGTAAAT 538
QY 737 TAAGTGTCTGAGTAATAATTTGGTAGCACAAACACTGAATGTACTAAATGTGCTC 796
Db 539 GCAGACTAACTTTTACTATAATGGTGGTTCCTTAAAGTGAAGCTCCTGGCGTTTAAAG 598
QY 797 CTAACTTTTACAATAATGCTCCTTAATTTCAATCCAGGTAAATAGTACATGCCTACCTT 856
Db 599 TTTTGTGCTGCTGCTGGCGTGCAGGTGTTGCTGCCGTTTACTAGTTAAATGTGTACCTT 658
QY 857 GCCAGCAAAATAAGATTATGGTCTGAAGCCACTCCAGGTGCTGCCGCTACTTTAGCCA 916
Db 659 GCCAACTAAACAAAACGATTCCTCT---GCCACTGCAGTGCCTTAAGCTAAATTTAGCCA 715
QY 917 AATAATGTAATTTGATGCGCTGATGGTACTGCAATTTGCTAGTGAGGACAACTAATATG 976
Db 716 CATATGCACTACTTAATGTCCAACCTGGCAGTGAATTCAGAGGAGTGACACTTGTT 775
QY 977 TAATATTATAACAGAAATGCTAAATTTGCTGCTCACTTTTATTTTGTGCTTAATTT 1036
Db 776 TTAGTATTCATCACAATAATGTTCTTAATGCAATGCTCAATTTTAAATGTTAATGTAATC 835
QY 1037 TCTAGGAGGAGTGTAGTGAAGCAAGCATGTCAGCAAAATAAGTTAAGCGCTGTAG 1096
Db 836 TCGAAGCAGGTGAAAGTTAATGTTAAAGTGTCCAGTAAGTAAACT---ACTCCAGCAC 892
QY 1097 CAACTGCAAGTGTGCTACTTTAATTCGATAATGTCCTTGAATGCCCTGCTGGTA 1156
Db 893 ATGCTCCAGGTAATGCTACTTAAGCCACATAATGTTTGACCAATGTCCTGCTGGTA 952
QY 1157 CTGACTCAGGAGTGAACATCTACTTATAAATAAGCAGCATCTGAATGTTTAAAT 1216
Db 953 CAGTACTTGATGAGGACATCACTAATTTGTAGCTCCGCACTGAATGTACTAAAT 1012
QY 1217 GTGCTGCCAACTTTTATACTACAAAATAACTGATGGGTAGCAGGTATGTACATGTA 1276
Db 1013 GTTCTGCTGGCTTTTTCATCAAAAACAACTGGTTTACAGGAGGTACTGTATACATGTA 1072
QY 1277 CTAGTTGTAATAAAAAATTAACCTTCTGGCGTGAAGCT 1314
Db 1073 CTGAATGTACTAAAAATAACTTCTGGTGCCACAGCT 1110
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RESULT 13
US-07-763-352A-14
; Sequence 14, Application US/07763352A
; GENERAL INFORMATION:
; APPLICANT: Clark, Theodore G.
; APPLICANT: Dickerson, Harry W.
; TITLE OF INVENTION: ICH IMMOBILIZATION ANTIGEN AND FISH
; TITLE OF INVENTION: VACCINE
; NUMBER OF SEQUENCES: 15
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Greenlee and Winner
; CITY: Boulder
; STATE: Colorado
; COUNTRY: USA
; ZIP: 80303
```

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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07763,352A
FILING DATE: 19910920
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Feiber, Donna M.
REGISTRATION NUMBER: 33,878
REFERENCE/DOCKET NUMBER: 15-91
TELECOMMUNICATION INFORMATION:
TELEPHONE: 303/499-8080
TELEFAX: 303/499-8089
TELEX: 823189
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 1936 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: cDNA to mRNA
FEATURE:
NAME/KEY: mat_peptide
LOCATION: 88..1269
FEATURE:
NAME/KEY: sig_peptide
LOCATION: 28..88
FEATURE:
NAME/KEY: CDS
LOCATION: 28..1272
OTHER INFORMATION: /codon= (seq: "taa", aa: Gln)
OTHER INFORMATION: /codon= (seq: "tag", aa: Gln)
US-07-763-352A-14
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Query Match 15.3%; Score 214.6; DB 3; Length 1936;
Best Local Similarity 55.4%; Pred. No. 6.2e-40;
Matches 591; Conservative 0; Mismatches 374; Indels 102; Gaps 5;

QY 344 GTGTTAATTTAGATAATTAATTTTATAATGAAATGCTCCAAATTTTAAATGAGGTGCTA 403
Db 194 GTGCTGCTTAAGGAGAAGCTAATGTTAATTAACCTTTTCGACGAAATAATGCTGTAGAG 253
QY 404 GTACATGCACAGCTTTCGGTAAACAGAGTTGTGTGTCATTTGCTGTAATGCGG 463
Db 254 GTATATGTGTACCAATGCAAAATAACAGAGTAGGCTCTGTTACCAATGAGGTGACTTAG 313
QY 464 CTACCATAGTCGCATAATGTAACGTCGCATGCTCTACTGTACTGCACCTTGATGGAG 523
Db 314 CTACTTAGCCACATANTGCAGTACTTAATGTCCTACTGGCACTGCATGATGATGGAG 373
QY 524 TAACTACTGATTATGTTAGATCATTACAGAAATGTTAATGATAGACTTAACCTTTTACT 583
Db 374 TGACAGATGTTTTTGATAGATCAGCCGCAATAATGTGTTAAATGCAAACTTAACCTTTTACT 433
QY 584 ATAATGGTAAATGTTAATTAATCTCTTCAATCCAGTAAAGTTAATGACACACCTTTGTC 643
Db 434 ATAATGGTGGTTCCTCTTAAGGTGAAGCTCCTGCCGTTTAAAGTTTTCGCTGCTGGTCTG 493
QY 644 CGGCAATTAACCTGCT-----AATG 664
Db 494 CGCGTTCAGGTTGTTGCTGCCGTTACTAGTTAATGCTACCTTGCCAACTAAACAAAACG 553
QY 665 TTGCTTTAAGCTACTTTTAGTAAATGCTACAATAACCGCATATATGTAAGCTTGCAATGCC 724
Db 554 ATTCTCCTGCCACTGCAGGTGCTTAAGCTAATTTAGCCACATAATGTAGCAATTAATGTC 613
QY 725 CTGATGCTACTATAAGTGTCTGCTGGAGT---AAATAATTTGGGTAGCACAAACACTGAAT 781
Db 614 CTACTGGCACTGTACTTGTATGATGGAGTGACACTGTTGTTTTTAAATACATCAGCCACATTAT 673
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Qy 782 GTACTAATTTGTCCTCTAAGTTTACAAATAATAATGCTCCTAATT----- 826
Db 674 GTGTTAAATGCAGACCTTAACCTTTTACTATTAATGGTGGTCTCTCTTAAGGTGAAGCTCCTG 733
Qy 827 -----TCAATCCAGTAAATACAT 847
Db 734 CGGTTTAAGTTTGTGCTGGTCTGCCGCTGCAGGCTGTGCTGCCGTTACTAGTTAAT 793
Qy 848 GCCTACCTTCCCGAGCAATAAGATTTATGGTCTGAAGCCACTGCAGGTGGTCCGCTA 907
Db 794 GTGTACCTTCCCAACTAACAACAAACGATCTCCT---GCCACTGCAGGTGCCCTAAGCTA 850
Qy 908 CTTTAGCCAAATTAATGTAATTTGTCATGCCCTGATGGTACTGCAATTTGCTAGTGAGCAA 967
Db 851 ATTTAGCCACATAATGCAGTACTTAATGTCCAACTGGCACTGCAATTTCAAGACGGAGTGA 910
Qy 968 CTAATTATGCTAATTTATAACACAGATGCTAAATTTGCTGCTAACTTTTATTTTATG 1027
Db 911 CACTTGTGTTTGTAAATTCATCCACATAATGTTCTTAATGCAATGCTAATTTACTTTT 970
Qy 1028 GTAATAATTTCTAGGCAGCAAGTAGTAGATGCAAGCAATGCTCCACCAATAAAGTTTAA 1087
Db 971 ATGGTAATCTCGAAGCAGTAAAGTTAATGTTTAAAGTGCCAGTAAGTAAACT---A 1027
Qy 1088 GCCTGTAGCAACTGCGAGTGGTACTGCTACTTTAATTTGCATAATGTCGCCCTTGAATGCC 1147
Db 1028 CTCAGCACATGCTCCAGGTAATGCTACTTAAAGCCACATAATGTTTGACCAATGTC 1087
Qy 1148 CTGCTGGTACTGCTACCCGATGAACACACATCTACTTTAATAAAGCAGCATCTGAAT 1207
Db 1088 CTGCTGGTACACTGCTGATGATGAACATCAACTTAATTTTGTAGCTTCCGCAACTGAAT 1147
Qy 1208 GTGTTAAATGCTGCCAACTTTTATACACAAAATAAAGTATGCTGGTACGAGTATG 1267
Db 1148 GTACTAAATGTTCTGCTGCTTTTGTGATCAAAAACACTGGTTTACAGCAGGACTG 1207
Qy 1268 ATACATGTAAGTGTAAATAAATAAATTAACCTCTGGCGCTGAAGCT 1314
Db 1208 ATACATGTAAGTGTAAATAAATAAATTAACCTCTGGTGGCACAGCT 1254

RESULT 14
US-09-196-161-1
; Sequence 1, Application US/09196161A
; GENERAL INFORMATION:
; APPLICANT: SIN, YOKE M
; APPLICANT: LAM, TOONG J
; APPLICANT: GONG, ZHIYUAN
; TITLE OF INVENTION: A RECOMBINANT VACCINE AGAINST FISH INFECTIOUS DISEASES
; FILE REFERENCE: RECOMBINANT VACCINE FOR FISH
; CURRENT APPLICATION NUMBER: US/09/196,161A
; CURRENT FILING DATE: 1998-11-20
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
; LENGTH: 316
; TYPE: DNA
; ORGANISM: Ichthyophthirius multifiliis
US-09-196-161-1

Query Match 7.9%; Score 110.4; DB 15; Length 316;
Best Local Similarity 65.3%; Pred. No. 1.5e-15;
Matches 162; Conservative 0; Mismatches 86; Indels 0; Gaps 0;

Qy 344 GTGTTAAATGTAGAAATTAATTTTATAATGAAATGCTCCAAATTTTAAATGCGAGGTGCTA 403
Db 2 GTGCTGCTTAAGGAGAAGCTAATGCTAATTAACCTTTTCGCAGCAAAATAATGCTGCTAGAG 61
Qy 404 GTACATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTTGACTGCTGGTAAATGCCG 463
Db 62 GTATATGTGTACCATGCCCAATAAACAGAGTAGGCTCTGTTACCAATGCAGGTGACTTAG 121
Qy 464 CTACCATAAGTCGCATTAATGTAACCTGCGCATGTCCTACTGGTACTGCACTTTGATGGAG 523
Db 122 CTACTTTAGCCACATAATGTCAGTACTTAAATGTCCTACTGGCACTGCACTTTGATGGAG 181
Qy 524 TAACACTGATTTAGTTAGTATTCATTCACAGAATGTGTTAAATGTAGACTTAACTTTTACT 583
Db 182 TGACAGATGTTTGTGATAGATCAGCCGCAATAATGTGTTAAATGCAAACTTAACTTTTACT 241
Qy 584 ATAATGGT 591
Db 242 ATAATGGT 249

Search completed: February 17, 2003, 01:35:53
Job time : 2281.72 secs
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Db 122 CTACTTTAGCCACATATGTCAGTACTTAATGCTCCTACTGGCACTGCACCTTGATGGAG 181
Qy 524 TAACACTGATTTAGTTAGTATTCACAAATGTGTTAAATGTAGACTTAACTTTTACT 583
Db 182 TGACAGATGTTTGTGATAGATCAGCCGCAATAATGTGTTAAATGCAAACTTAACTTTTACT 241
Qy 584 ATAATGGT 591
Db 242 ATAATGGT 249

RESULT 15
US-09-196-161-2
; Sequence 2, Application US/09196161A
; GENERAL INFORMATION:
; APPLICANT: SIN, YOKE M
; APPLICANT: LAM, TOONG J
; APPLICANT: GONG, ZHIYUAN
; TITLE OF INVENTION: A RECOMBINANT VACCINE AGAINST FISH INFECTIOUS DISEASES
; FILE REFERENCE: RECOMBINANT VACCINE FOR FISH
; CURRENT APPLICATION NUMBER: US/09/196,161A
; CURRENT FILING DATE: 1998-11-20
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 2
; LENGTH: 316
; TYPE: DNA
; ORGANISM: Ichthyophthirius multifiliis
US-09-196-161-2

Query Match 7.9%; Score 110.4; DB 15; Length 316;
Best Local Similarity 65.3%; Pred. No. 1.5e-15;
Matches 162; Conservative 0; Mismatches 86; Indels 0; Gaps 0;

Qy 344 GTGTTAAATGTAGAAATTAATTTTATAATGAAATGCTCCAAATTTTAAATGCGAGGTGCTA 403
Db 2 GTGCTGCTTAAGGAGAAGCTAATGCTAATTAACCTTTTCGCAGCAAAATAATGCTGCTAGAG 61
Qy 404 GTACATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTTGACTGCTGGTAAATGCCG 463
Db 62 GTATATGTGTACCATGCCCAATAAACAGAGTAGGCTCTGTTACCAATGCAGGTGACTTAG 121
Qy 464 CTACCATAAGTCGCATTAATGTAACCTGCGCATGTCCTACTGGTACTGCACTTTGATGGAG 523
Db 122 CTACTTTAGCCACATAATGTCAGTACTTAAATGTCCTACTGGCACTGCACTTTGATGGAG 181
Qy 524 TAACACTGATTTAGTTAGTATTCATTCACAGAATGTGTTAAATGTAGACTTAACTTTTACT 583
Db 182 TGACAGATGTTTGTGATAGATCAGCCGCAATAATGTGTTAAATGCAAACTTAACTTTTACT 241
Qy 584 ATAATGGT 591
Db 242 ATAATGGT 249

Search completed: February 17, 2003, 01:35:53
Job time : 2281.72 secs
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